

TABLE III

Relative Position									
	1	2	3	4	5	6	7	8	9
DRB1*0101	Y* (47%)	X	X	M (48%)	X	A (34%)	X	X	L (43%)
	F (26%)			L (28%)		G (23%)			M (13%)
						S (09%)			A (13%)
DRB1*0401	W (43%)	X	X	M (23%)	X	T (60%)	L (34%)	X	X
	Y (31%)			A (19%)		S (12%)	Q (20%)		
				V (13%)			M (10%)		
				L (12%)			N (10%)		
DRB1*1101	W (67%)	X	X	M (33%)	X	R (61%)	X	X	X
				L (23%)		K (12%)			
				V (13%)					

IN THE CLAIMS:

Please amend the Claims as follows without prejudice.

46. A peptide derived from glutamic acid decarboxylase having a length of at most 25 amino acids and comprising

(a) a peptide of at least 6 amino acids of an amino acid sequence selected from the group consisting of SEQ ID NOS: 2, 3 and 20-39, or

(b) a peptide or peptide derivative having a length of 6 to 25 amino acids which exhibits a specificity or/and affinity which is equivalent to that of the peptide (a) and